## Genopsys.708.ST25

## SEQUENCE LISTING

- × × ×

```
<110> Lietz, Eric
  <120> RANDOM TRUNCATION AND AMPLIFICATION OF NUCLEIC ACID
  <130> 22477-708
  <150> US C9/518,335
  <151> 2000-03-03
  <150> PCT/US01/07016
  <151> 2001-03-05
  <160> 2
<170> PatentIn version 3.1
43
<210> 1
<211> 21
¥7 <220>
4 <223> 5'-Primer
Pi <220>
C <221> misc feature
(223> "n" residues in position 13-21 represent adenosine, thymidine, g
(3 u
        anosine, cytidine, uridine or inosine.
Pet.
  <400> 1
  aaaatacata tgnnnnnnnn n
                                                                  2
  <210> 2
  <211> 31
  <212> DNA
  <213> Artificial Sequence
  <220>
  <223> 3-Primer
  <220>
  <221> misc_feature
  <222> (23)..(31)
  <223> "n" residues in position 23-31 represent adenosine, thymidine, q
```

Genopsys.708.ST25

anosine, cytidine, uridine or inosine.

<400> 2 ataagtgctt cactactaat tannnnnnn n

3